CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/912,628

ENTERED

CRF Processing Date: 9/12/2001
Edited by: _____
Verified by: ___AL___ (STIC sta[

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section. specifically:
_____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings). specifically:
_____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: ·. .
_____

☐ Deleted extra, invalid, headings used by an applicant. specifically:
_____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response. specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field. specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☑ Other: *inserted closing bracket in <110>*
_____
_____

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.
3/1/95

```
          RAW SEQUENCE LISTING                DATE: 09/12/2001
          PATENT APPLICATION:  US/09/912,628  TIME: 11:52:23

            Input Set : A:\Pto.amc
            Output Set: N:\CRF3\09122001\I912628.raw
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        2 <110> APPLICANT: Ni et al.
        4 <120> TITLE OF INVENTION: Human Serpin Polynucleotides, Polypeptides, and Antibodies
        6 <130> FILE REFERENCE: PT001P2
C-->    8 <140> CURRENT APPLICATION NUMBER: US/09/912,628
        9 <141> CURRENT FILING DATE: 2001-07-26
       11 <150> PRIOR APPLICATION NUMBER: PCT/US01/02484
       12 <151> PRIOR FILING DATE: 2001-01-26
       14 <150> PRIOR APPLICATION NUMBER: 60/178,769
       15 <151> PRIOR FILING DATE: 2000-01-28
       17 <150> PRIOR APPLICATION NUMBER: PCT/US00/05082
       18 <151> PRIOR FILING DATE: 2000-02-29
       20 <160> NUMBER OF SEQ ID NOS: 17
       22 <170> SOFTWARE: PatentIn Ver. 2.0
       25 <210> SEQ ID NO: 1
       26 <211> LENGTH: 733
       27 <212> TYPE: DNA
       28 <213> ORGANISM: Homo sapiens
       30 <400> SEQUENCE: 1
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       33 tctcccggac tcctgaggtc acatgcgtgg tggtggacgt aagccacgaa gaccctgagg   180
       34 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg   240
       35 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact   300
       36 ggctgaatgg caaggagtac aagtgcaagg tctccaacaa agccctccca accccatcg    360
       37 agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc   420
       38 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct   480
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       41 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc   660
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       54 aatctactgt gtgtccccgg ccaatgcccc cagtgcatac ccccgccctt cctccacaaa   180
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67 caagatgagg caactggaac aggccttgtc agccagaaca ctgagaaagt ggagccactc      960
68 actccagaaa aggtggatag aggtgttcat ccccagattt tccatttctg cctcctacaa     1020
69 tctggaaacc atcctcccga agatgggcat ccaaaatgtc tttgacaaaa atgctgattt     1080
70 ttctggaatt gcaaagagag actccctgca ggtttctaaa gcaacccaca aggctgtgct     1140
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77 acacccatta accccagtca gtgccctttt ccacaaattc tcccaggtaa ctagcttcat     1560
78 gggatgttgc tgggttacca tatttccatt ccttggggct cccaggaatg gaaatacgcc     1620
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 91 gtcaaagaga ctttcttcaa tttatccaag aggtattttg atacagagtg cgtgcctatg     180
 92 aattttcgca atgcctcaca ggccaaaagg ctcatgaatc attacattaa caaagagact     240
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 94 gattacatct tgttcaaagg gaaatggttg accccatttg accctgtctt caccgaagtc     360
 95 gacactttcc acctggacaa gtacaagacc attaaggtgc ccatgatgta cagtgcaggc     420
 96 aagtttgcct ccacctttga caagaatttt cgttgtcatg tcctcaaact gccctaccaa     480
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 98 gactacctga ccacagactt ggtggagaca tggctcagaa acatgaaaac cagaaacatg     600
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104 tttctgggca gggtggtgaa tccgactctc ctataattca ggacacgcat aagcacttcg     960
105 tgctgtagta gatgctgaat ctgaggtatc aaacacacac aggataccag caatggatgg    1020
106 caggggagag tgttcctttt gttcttaact agtttagggt gttctcaaat aaatacagta    1080
107 gtccccactt atctgagggg gatacattca aagacccca gcagatgcct gaaacggtgg    1140
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109 ttataaatta ggcacagtaa gagattaaca ataataacaa cattaagtaa aatgagttac    1260
110 ttgaacgcaa gcactgcaat accataacag tcaaactgat tatagagaag gctactaagt    1320
111 gactcatggg cgaggagcat agacagtgtg gagacattgg gcaaggggag aattcacatc    1380
112 ctgggtggga cagagcagga caatgcaaga ttccatccca ctactcagaa tggcatgctg    1440
113 cttaagactt ttagattgtt tatttctgga atttttcatt taatgttttt ggaccatggt    1500
114 tgaccatggt taactgagac tgcagaaagc aaaaccatgg ataagggagg actactacaa    1560
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116 aaaaaaaaaa aa                                                         1632
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121 <212> TYPE: DNA
122 <213> ORGANISM: Homo sapiens
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125 <221> NAME/KEY: SITE
126 <222> LOCATION: (635)
127 <223> OTHER INFORMATION: n equals a,t,g, or c
129 <220> FEATURE:
130 <221> NAME/KEY: SITE
131 <222> LOCATION: (655)
132 <223> OTHER INFORMATION: n equals a,t,g, or c
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137 tcaaacaact tatttttttc cccatgagca tatcatcagc cttggccatg gtttcatgg      180
138 gggcaaaggg aaacactgca gctcagatgt ctcaggcact ttgttttagt aaaatcggag      240
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140 acactgaata tgtgcttaga actgccaacg ggctctttgg agaaaagtct tatgatttcc      360
141 tcacaggttt tacagattcc tgtggcaaat tctaccaagc aacgataaaa cagctagact      420
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144 cctcttgtta ctgcaaagcc tgcctttcac agcccctact ggttcactct attcccaaat      600
145 gcaactctcc tgtgaccccg catggcatgt ggtgnccctcc ctccctgtga gcagntgtga      660
146 ctaataaact gccgccaatt tcatctgtaa aaaaaaaaaa aaaaaa                      706
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150 <211> LENGTH: 435
151 <212> TYPE: PRT
152 <213> ORGANISM: Homo sapiens
154 <400> SEQUENCE: 5
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159              20                  25                  30
161 Cys Ala Pro Ile Tyr Cys Val Ser Pro Ala Asn Ala Pro Ser Ala Tyr
162          35                  40                  45
164 Pro Arg Pro Ser Ser Thr Lys Ser Thr Pro Ala Ser Gln Val Tyr Ser
165      50                  55                  60
167 Leu Asn Thr Asp Phe Ala Phe Arg Leu Tyr Arg Arg Leu Val Leu Glu
168  65                  70                  75                  80
170 Thr Pro Ser Gln Asn Ile Phe Phe Ser Pro Val Ser Val Ser Thr Ser
171              85                  90                  95
173 Leu Ala Met Leu Ser Leu Gly Ala His Ser Val Thr Lys Thr Gln Ile
174             100                 105                 110
176 Leu Gln Gly Leu Gly Phe Asn Leu Thr His Thr Pro Glu Ser Ala Ile
177             115                 120                 125
179 His Gln Gly Phe Gln His Leu Val His Ser Leu Thr Val Pro Ser Lys
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W--> 145

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180        130                    135                   140
182 Asp Leu Thr Leu Lys Met Gly Ser Ala Leu Phe Val Lys Lys Glu Leu
183 145                    150                  155                 160
185 Gln Leu Gln Ala Asn Phe Leu Gly Asn Val Lys Arg Leu Tyr Glu Ala
186                    165                  170                 175
188 Glu Val Phe Ser Thr Asp Phe Ser Asn Pro Ser Ile Ala Gln Ala Arg
189                180                  185                 190
191 Ile Asn Ser His Val Lys Lys Lys Thr Gln Gly Lys Val Val Asp Ile
192            195                  200                 205
194 Ile Gln Gly Leu Asp Leu Leu Thr Ala Met Val Leu Val Asn His Ile
195     210                  215                 220
197 Phe Phe Lys Ala Lys Trp Glu Lys Pro Phe His Pro Glu Tyr Thr Arg
198 225                    230                  235                 240
200 Lys Asn Phe Pro Phe Leu Val Gly Glu Gln Val Thr Val His Val Pro
201                245                  250                 255
203 Met Met His Gln Lys Glu Gln Phe Ala Phe Gly Val Asp Thr Glu Leu
204            260                  265                 270
206 Asn Cys Phe Val Leu Gln Met Asp Tyr Lys Gly Asp Ala Val Ala Phe
207        275                  280                 285
209 Phe Val Leu Pro Ser Lys Gly Lys Met Arg Gln Leu Glu Gln Ala Leu
210     290                  295                 300
212 Ser Ala Arg Thr Leu Arg Lys Trp Ser His Ser Leu Gln Lys Arg Trp
213 305                    310                  315                 320
215 Ile Glu Val Phe Ile Pro Arg Phe Ser Ile Ser Ala Ser Tyr Asn Leu
216                325                  330                 335
218 Glu Thr Ile Leu Pro Lys Met Gly Ile Gln Asn Val Phe Asp Lys Asn
219            340                  345                 350
221 Ala Asp Phe Ser Gly Ile Ala Lys Arg Asp Ser Leu Gln Val Ser Lys
222        355                  360                 365
224 Ala Thr His Lys Ala Val Leu Asp Val Ser Glu Glu Gly Thr Glu Ala
225     370                  375                 380
227 Thr Ala Ala Thr Thr Thr Lys Phe Ile Val Arg Ser Lys Asp Gly Pro
228 385                    390                  395                 400
230 Ser Tyr Phe Thr Val Ser Phe Asn Arg Thr Phe Leu Met Met Ile Thr
231                405                  410                 415
233 Asn Lys Ala Thr Asp Gly Ile Leu Phe Leu Gly Lys Val Glu Asn Pro
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236 Thr Lys Ser
237         435
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241 <211> LENGTH: 311
242 <212> TYPE: PRT
243 <213> ORGANISM: Homo sapiens
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252 Ala Phe Ile His Lys Asp Phe Asp Val Lys Glu Thr Phe Phe Asn Leu
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255 Ser Lys Arg Tyr Phe Asp Thr Glu Cys Val Pro Met Asn Phe Arg Asn
256       50                      55                  60
258 Ala Ser Gln Ala Lys Arg Leu Met Asn His Tyr Ile Asn Lys Glu Thr
259 65                      70                  75                  80
261 Arg Gly Lys Ile Pro Lys Leu Phe Asp Glu Ile Asn Pro Glu Thr Lys
262               85                      90                  95
264 Leu Ile Leu Val Asp Tyr Ile Leu Phe Lys Gly Lys Trp Leu Thr Pro
265          100                     105                 110
267 Phe Asp Pro Val Phe Thr Glu Val Asp Thr Phe His Leu Asp Lys Tyr
268      115                     120                 125
270 Lys Thr Ile Lys Val Pro Met Met Tyr Ser Ala Gly Lys Phe Ala Ser
271      130                     135                 140
273 Thr Phe Asp Lys Asn Phe Arg Cys His Val Leu Lys Leu Pro Tyr Gln
274 145                     150                 155                 160
276 Gly Asn Ala Thr Met Leu Val Val Leu Met Glu Lys Met Gly Asp His
277               165                     170                 175
279 Leu Ala Leu Glu Asp Tyr Leu Thr Thr Asp Leu Val Glu Thr Trp Leu
280          180                     185                 190
282 Arg Asn Met Lys Thr Arg Asn Met Glu Val Phe Phe Pro Lys Phe Lys
283      195                     200                 205
285 Leu Asp Gln Lys Tyr Glu Met His Glu Leu Leu Arg Gln Met Gly Ile
286      210                     215                 220
288 Arg Arg Ile Phe Ser Pro Phe Ala Asp Leu Ser Glu Leu Ser Ala Thr
289 225                     230                 235                 240
291 Gly Arg Asn Leu Gln Val Ser Arg Val Leu Gln Arg Thr Val Ile Glu
292               245                     250                 255
294 Val Asp Glu Arg Gly Thr Glu Ala Val Ala Gly Ile Leu Ser Glu Ile
295          260                     265                 270
297 Thr Ala Tyr Ser Met Pro Pro Val Ile Lys Val Asp Arg Pro Phe His
298      275                     280                 285
300 Phe Met Ile Tyr Glu Glu Thr Ser Gly Met Leu Leu Phe Leu Gly Arg
301      290                     295                 300
303 Val Val Asn Pro Thr Leu Leu
304 305                     310
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307 <210> SEQ ID NO: 7
308 <211> LENGTH: 215
309 <212> TYPE: PRT
310 <213> ORGANISM: Homo sapiens
312 <220> FEATURE:
313 <221> NAME/KEY: SITE
314 <222> LOCATION: (211)
315 <223> OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids

317 <400> SEQUENCE: 7

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321 His Gly Cys Ser Ile Arg Ser Lys Trp His Ile Cys Ile Lys Pro Phe
322               20                  25                  30
324 Glu Lys Ala Arg Gly Lys Gln Leu Lys Gln Leu Ile Phe Phe Pro Met
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9/12/01

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/912,628

DATE: 09/12/2001
TIME: 11:52:24

Input Set : A:\Pto.amc
Output Set: N:\CRF3\09122001\I912628.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application Number
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:385 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:388 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:441 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:444 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:537 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
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L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17

9/12/01